

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoretic
Q5F2E7	NUFP2_MOUSE	Mus musculus	FMR1-interacting protein NUFIP2	52.077887	T369;S376;S380;S429;T533	T88;S113;S114;S213;S215;T219;T220;T221;S304;S376;T569;S570;S589;S605;S626;T630;S634;S649;S652;S689	22517741;22645316;34887587;40885482;36852467;40997131	MEEKPGQPQPOHHHSHHHPHHHP QQQQQQSHHHHHYYFYNHSHNH HHHHHHQPHQYQLQHAEGSPKA QPKPLKHEQKHTLQQHQETPKKKT GYGEINGNAGEREISLKSLSDEATN PISRVLNGNQVVETSLKQTVKTSTF GKAGIKTKNFIQKNSMDKKNGKSYE NKSGETQAVDKTDTIAIPNGVITSSS GYITNGYMSKGADNDGSGSESGYTT PKKRKARRNSAKGCENLNLVQDKI MQETSVPAKQGLETLKPDYSEQKG MRVDGSKPIWKYETGPGGTSRGPKA MGDVLRSDDIKFGLSSKKFDDRPK GKHASAAASKEDSWTLFKPPVFPV DNSSAKIVPKISYASKVKENLNKTVO NSSVSPSSSSSSSTGETQTQSSSRL SQVPMALKSVTSASFNGPVLGAT DGSVYPSGGQPLLTTAANTLTPISTG TDSVLQDMSLASAAVEQIKSSLFIYP SNMQTVLLSAQVDLPSQTDQQNLG DIFQNWGLSFINEPSAGPETVIGKS SDHKVMEVTFQGEYPATLVSQGAII PSGTEHPVFPKAYELEKRTSPQVLG HILKPGTTESGALS LDP SHIGDLQKA DTSSQGALVFLSKDYEIENQNPLASP TNTLLGSAKEQRYQRGLERNDSWG SFDLRAAIVYHTKEMESIWNLQKQD PKRIITYNEAMDSPDQ	True	False	3.327	4.172	0.791	0.741